

Serial No.: 10/806,899

IN THE SPECIFICATION

Please replace the Sequence Listing with the Substitute Sequence Listing submitted herewith. No new matter has been added.

Please replace Table 1 on page 43 with the following replacement Table 1:

TABLE 1

Primer Sequences Used for dHPLC Assay Analysis of SCN1A

Exon	Forward Primer	Reverse Primer	Size (bp)
1	CCTCTAGCTCATGTTTTCATGAC (SEQ ID NO: 59)	TGCAGTAGGCAATTAGCAGC (SEQ ID NO: 60)	448
2	CTAATTAAGAAGAGATCCAGTGACAG (SEQ ID NO: 61)	GCTATAAGTGTTCACAGATCATGTAC (SEQ ID NO: 62)	356
3	CCCTGAATTTTGGCTAAGCTGCAG (SEQ ID NO: 63)	CTACATTAAGACACAGTTTCAAAATCC (SEQ ID NO: 64)	263
4	GGGTACGTTTCAATTGTATG (SEQ ID NO: 65)	GCAACCTATTCTTAAAGCATAAAGACTG (SEQ ID NO: 66)	358
5	AGGCTCTTTGTACCTACAGC (SEQ ID NO: 67)	CATGTAGGGTCCGTCCTCATT (SEQ ID NO: 68)	200
6	CACACGTGTTAAGTCTTCATAGT (SEQ ID NO: 69)	AGCCCTCAAGTATTTATCCT (SEQ ID NO: 70)	394
7	GAACCTGACCTTCCTGTTCTC (SEQ ID NO: 71)	GTTGGCTGTTATCTTCAGTTTC (SEQ ID NO: 72)	241
8	AAAGGCAGCAGAACGACTTG (SEQ ID NO: 73)	GGATAGAGAACTCAAGTCTC (SEQ ID NO: 74)	322
9	TTGAAAAGTTGAAAGCCACCAC (SEQ ID NO: 75)	CCACCTGCTCTTAGGTACTC (SEQ ID NO: 76)	363
10	GCCATGCAATACTTCAGCCC (SEQ ID NO: 77)	CACAACAGTGGTTGATTCAGTTG (SEQ ID NO: 78)	480
11(1)	TGAATGCTGAAATCTCCTTCTAC (SEQ ID NO: 79)	CTCAGGTTGCTGTTGCGTCTC (SEQ ID NO: 82)	306
11(2)	GATAACGAGAGCCGTAGAGAT (SEQ ID NO: 81)	TCTGTAGAAACACTGGCTGG (SEQ ID NO: 82)	315
12	CATGAAAATTCACCTGTGCACC (SEQ ID NO: 83)	CAGCTCTTGAATTAGACTGTC (SEQ ID NO: 84)	347
13	ATCCTTGGGAGGTTTAGAGT (SEQ ID NO: 85)	GCATGAAGGATGTTGAAAG (SEQ ID NO: 86)	510
14	CATTGTGGGAAAATAGCATAAGC (SEQ ID NO: 87)	GCTATGCAGAACCCCTGATTG (SEQ ID NO: 88)	339
15(1)	TGAGACGGTTAGGCAGATC (SEQ ID NO: 89)	AGAAATCATTCATGTGCCAGC (SEQ ID NO: 90)	348
15(2)	GTCCTGGCCATCATCGTCTTC (SEQ ID NO: 91)	ACATGTGCACAAATGTGCAGG (SEQ ID NO: 92)	350
16(1)	GTGTTGTTCTTCTCATCAAG (SEQ ID NO: 93)	CACCTGCTGCCAGTTCCTATAC (SEQ ID NO: 94)	458
16(2)	CAACAGTCTTCATTAGGAAAC (SEQ ID NO: 95)	ACCTTCCCACACCTATAGAAATC (SEQ ID NO: 96)	353
17	CTTGGCAGGCAACTTATTACC (SEQ ID NO: 97)	CAAGCTGCACTCCAAATGAAAG (SEQ ID NO: 98)	232
18	TGGAAGCAGAGACACTTATCTAC (SEQ ID NO: 99)	GTGCTGTATCACCTTTTCTTAATC (SEQ ID NO: 100)	234
19	CCTATTCCAATGAAATGTCATATG (SEQ ID NO: 101)	CAAGCTACCTTGAAACAGAGAC (SEQ ID NO: 102)	318
20	CTACACATTGAATGATGATTCTGT (SEQ ID NO: 103)	GCTATATACAATACTTCAGGTTCT (SEQ ID NO: 104)	216
21	ACCAGAGATTACTAGGGGAAT (SEQ ID NO: 105)	CTGGGCTCATAACTTGTAATAAC (SEQ ID NO: 106)	513
22	ACTGCTTGGTCCAAATCTG (SEQ ID NO: 107)	TTTGATTAAATTTTACCACCTGATC (SEQ ID NO: 108)	267
23	AGCACCAAGTACATTTCCAAAC (SEQ ID NO: 109)	GGCAGAGAAAAACACTCCAAAG (SEQ ID NO: 110)	271
24	GACACAGTTTAAACCAGTTTG (SEQ ID NO: 111)	TGTGAGACAAGCATGCAAGTT (SEQ ID NO: 112)	207
25	CAGGGCCAAATGACTACTTTGCTG (SEQ ID NO: 113)	CTGATTGCTGGGATGATCTTGAATC (SEQ ID NO: 114)	477
26(1)	CAGGACTCTGAACCTTACCTTG (SEQ ID NO: 115)	ATTCCAACAGATGGGTCCCA (SEQ ID NO: 116)	534
26(2)	TCCTGCGTTGTTAAACATCGG (SEQ ID NO: 117)	AGGCAGCTGCAAACTGAGAT (SEQ ID NO: 118)	504
26(3)	TGGAAGCTCAGTTAAGGGAGA (SEQ ID NO: 119)	GTAGTGATTGGCTGATAGGAG (SEQ ID NO: 120)	480
26(4)	CCGATGCAACTCAGTTCAATGGA (SEQ ID NO: 121)	TGCCTTCTTGCTCATGTTTTCACA (SEQ ID NO: 122)	555
26(5)	AGAGCGATTTCATGGCTTCCAAATCC (SEQ ID NO: 123)	TGCTGACAAAGGGGTCACTGTCT (SEQ ID NO: 124)	526

Note: Primer sequences are listed 5' to 3'. Due to the large size of exons 11, 15, 16, and 26, the exons were split into two or more overlapping amplicons.